

***nmt1* promoter**

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-1178 TGATCAGAA AATTATCGCC ATAAAAGACA GAATAAGTCA TCAGCGGTTG TTTCATTTC TA
-1116 TATTTTTTTT TTATTTTTTT ATTTTTTAAT AAGGGAAAAT TTAACGTCTA AGGATACAGA AG
-1054 ATTGTTAGCA CATTAAAGTA ATAAAGGCTT AAGTAGTAAG TGCCTTAGCA TGTATTGTA TT
-992 TCAAAGGACA TAATCTAAAA TAATAACAAT ATCATTCTC ACAAGTTATT CAATTTTCTT TT
-930 TTTTTTCTAA TAATATCAAG AATGTATTAT TTGTTTGACA TAAGTCAACT AATTATTTA AT
-868 ATGCTGGATT AATCTTGCAG ACATGTAAAT TAACAAGTTT TAGTCAAATA ACGTTGAAGT TT
-806 CAATGAACTC AAATAATTTC TCTTTTTTTT TATATAACCA TATGTCTAAT CTGATTTATA TT
-744 TTCGCAGGA TCAACTGAAG TTATGACATT TGGATTGGAT CACTTATAAC CTTGGTCGCC AA
-682 ATAATACAAA AATCAGCGTT ATAAAACAAA GAAGGTTTTT GTTAAGAAAT TAATCCTCTT TC
-620 TTGATAAGAA AGTTGAACCG AAATTGCAGA TACTGATATA TGAAAAAAT ACCCACAATT TT
-558 GGAATAGCG CAAGCCTCAA TTAAACAAT AGGTGAGGAC ACATGATAAT GACCTCAATG AT
-496 TGTAGAAGA AAAGAGCCTC ATTACAAAAT CGAAAAATGA ATGGTTGGGT ACAAGTTTCC AA
-434 AACATGGTAA AGTGGACTTT GCGTATGAGA CGTAAATAGA AAAAAACACT TGTATATGT TT
-372 TCTAGAATTA TTGTTGTCTC TTTATGGTTG GATGATGCAA AATAGTAATT TCGGTTAGTT GC
-310 TGTA AACAC CACGAGACAA ATAGATATGG ATATTATTA AATCAGGAAA AACGTAATC TC
-248 GGCTACTGGA TGGTTCAGTC ACCCAACGAT TACTGGGAG AGAAAACAGG GCAAAAGCAA AG
-186 CTTAAAGGAA TCCGATTGTC ATTCGGCAAT GTGCAGCGAA ACTAAAAACC GGATAATGGA CC
-124 TGTTAATCGA AACATTGAAGA TATATAAG GAAGAGGAAT CCTGGCATAT CATCAATTGA AT
-62 AAGTTGAATT AATTATTTC AATCTATTCT CACTTTCTGA CTTATAGTC CTTTGTAAA TC
1 ATGTCTACTA ACAAGATCAC TT

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SEQ ID No.1 : *nmt-185* promoter

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AAAGGAATCCGATTGTCATTTCGGCAATGTGCAGCGAAACTAAAAACCGGATAATGGAGC
TGTTAATCGAAACATTGAAGATATATAAGGAAGAGGAATCCTGGCATATCATCAATTGA
ATAAGTTGAATTAATTATTTCAATCTCATTCTCACTTTCTGACTTATAGTCGCTTTGTTAA
ATCAT

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SEQ ID No.2 : *nmt-146* promoter

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TAAAAACCGGATAATGGACCTGTTAATCGAAACATTGAAGATATATAAGGAAGAGGAATCCTG
GCATATCATCAATTGAATAAGTTGAATTAATTATTTCAATCTCATTCTCACTTTCTGACTTATAGTC
GCTTTGTTAAATCAT

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Figure 1.